FIG. 1 (SEQ. ID NO: 1 & 2)

	8	180	270	360	450	5 <u>4</u> 0	630	22	810	8	86	1080	1147
40 30 234567890 1234567800 123456780 123456780 1234567800 1234567800 1234567800 12345678000 12345678000000000000000000000	1234567890 1234567890 1234567899 1234567899 1234567890 123456780 1234567890 123456780 12345780 123456780 123456780 123456780 123456780 123456780 123456780 123456780 123456780 123456780 123456780 123456780 123456780 123456780 123456780 123456780 123456780 123456780 123456780 123456780	TOTATIONAS ANCHITICAS ANACTITICO TOCCICIONAT COTOCOANTA GITTICOTOS TICCACITOS AGONATICO CIR E D V R E F A K V F L P V F L T I V P V I G L A G N S	AUGSDASTOS CAATTATOS CIATTACAAG AAACAGAGAA CCAAAACAGA TGIGTACAGC CICAATTIGS CUSDACAGA TTTACTOST ${ m M}$ V V A I Y A Y Y K K Q R T K T D V Y I L N L A V A D L L L	CENTICACIC TOCCITITIG COCTOTIVATE CENCITICAGE COTCOCCITITIN AGASSAAATA AUGICEAAA TAACITEAGE CITIGIACAA. L $oldsymbol{r}$ is a $oldsymbol{r}$ in	CIRARCITIC TCICICCAR COGITICIO CCTICIBAICA COMPACIOCA GIRACIRARCA TCCCCAGOCA ATCACARGIG $_{ m L}$ N F V S G M Q F L A C I S I D R Y V A V T K V P S Q S G V	GRADA CHA CHARACHA CHARACHA CHOCHARACHA CHARACHA CHARACHA TITITINIA RAIDAHACHA CHARACHA CHARA	AMICERCET GENETICE TITE THE TERM THE SHEAR ACCUTIONITY CARMICETIC ACMITTION IN A R C I P I F P R Y L G T S M K A L I Q M L E I C I G F V	GINDOCTITIC TINITATIONS GITGIOCTINC TITINICAGG CANGGACACT CANGAAGATIS AAMINICOG ACCCINADA V P F L I M G V C Y F I T A R T L M K M P N I K I S R P L K	GITCHCOLOR CAGNOTIME AGENTICAL GROWING THE COMPONENT CONTROL OF Γ of	CIGNICACIA COTOCAACIC ATCOPCATOS COMOCAACIT CACAGAACOC ATCOCACICIT TICACOCTIC COTCAACOCA $f L$ I I I S C N M S K R M D I A I Q V I E S I A L F H S C L N P	ATCCTTIATG TITTIATICGS ACCULCITIC ARAPACINGS TIMIGAANGT GOCCAAGANA TATGOSTOCT GOGGANACA GAGAANATAT IN TANA YU MKU AKKYG SWRR QRQ RQS	GICCHOCHET TICCITITION TICTIONGED CONTROPORTING TITTINGCRUT TRANGSTRAN ACTIGNIC CITTIGNIG VEEFPPFOSEGPTEPTCL	CATECATATIC AATGESTECTT TOCCCICAAA TAAAACATCT GOOTIDGTICT GAAAAAAAA AAAAAAA DIYE CPPLK NICLILKKKKK
30	UKBLUCELLI UKBURU UKBLUCELLI UKBLUCELLI UKBLUCELLI UKBURU UKBLUCELLI UKBLUCELLI UKBURU UKBURU UKBUR	e f a k v f	TIMITIACAAG AAACAGAGA Y Y K K Q R	XETICHTRAFF GEAGFICEN A V N A V H	CACTUCIES COTICIENTO QUE LE A C. I.	CRC V W M A	FPRYLG	VCY FIT	ACTITIONIT GROACION V P. I. V T. Q.	ECCHANCIC ATGERCHI S K R M D I	ACCULTIC ARAPACTS A S F K N Y	TOTAGEST COTACTOR	DECEMBARY TRANSCRE
20	1234567890 L 2 N Q 3 N Q	ANCAUTOTOMO AN D V R 1	CANTITATION C	TOCTITITIE O	TOTOTOGRAPH O	CCIGGACOR C	CONTROCOR I	TIMITATOS C	COCIONIBLE P	A COTTOCAACAT G	S TITITIBATOOG A	r miccumica 1	S AAGGAGGTT 1
10	1234567890 ATCCCTTTCC M A L E	TOTATIONARG C I K B	ATTOSTINGE M V V A	CENTICACIC L F T L	CIPAACTITIC L N F V	GENERACE C	AMICCIPACST N A R C	CIPOCCITIC V P F L	GITCICCICA V L L T	CICATORICA LITS	MOCTIFIATG I L Y V	GTCCPCCPGT V E E F	CKTPCATATG D T Y E

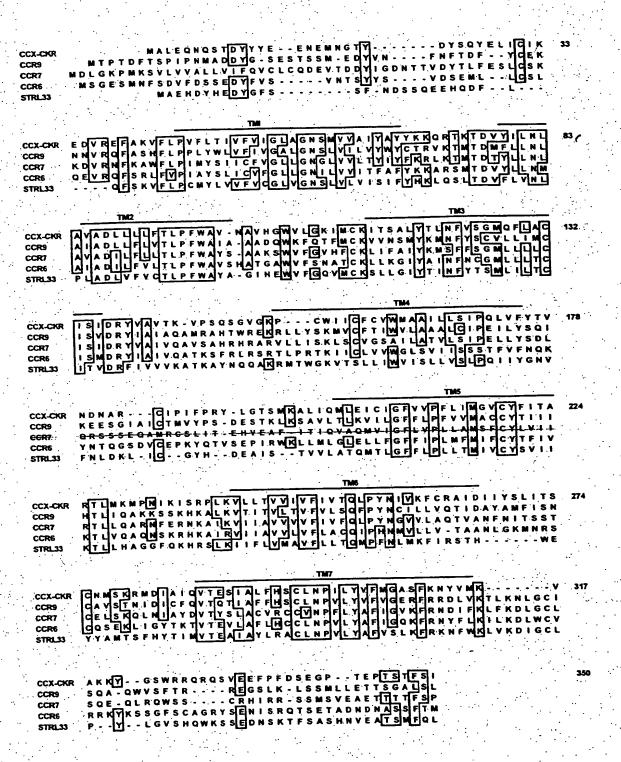


FIG. 2(a)

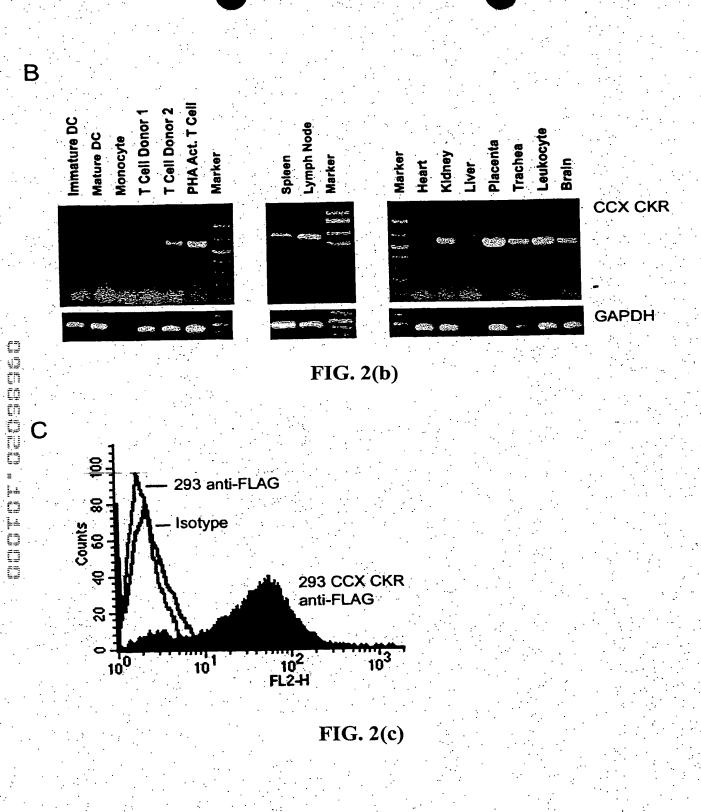


FIG. 3(a)

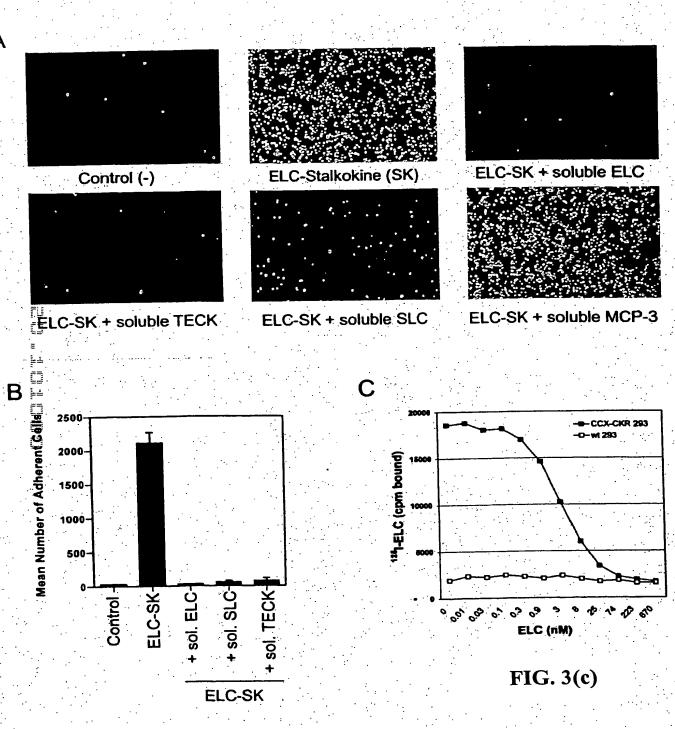


FIG. 3(b)

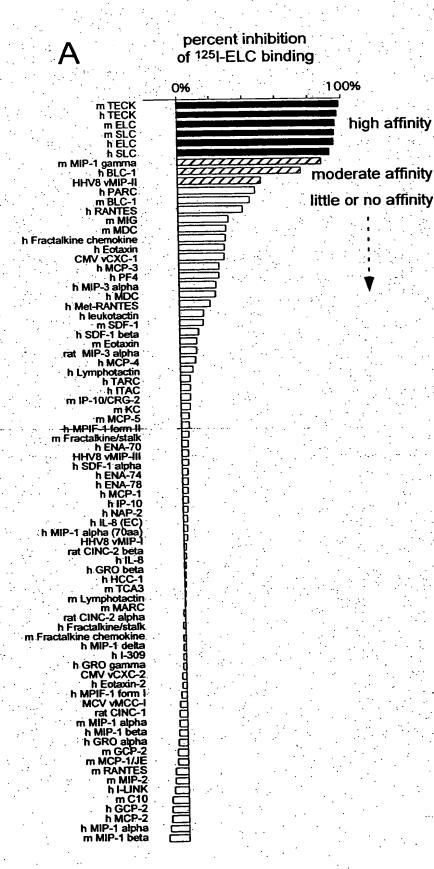
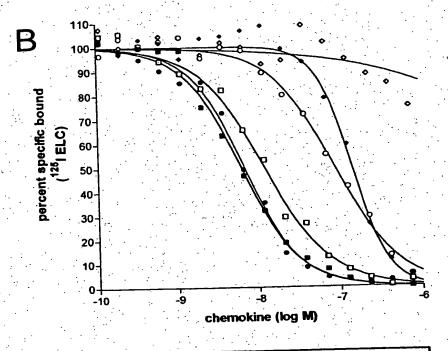


FIG. 4(a)



human chemokines	murine che	murine chemokines			
IC-50 ■ h ELC 6 nM □ h SLC 12 nM • h TECK 7 nM • h BLC-1 140 nM • h HHV8 vMIP-II 90 nM • h MCP-3 >2000 nM	m ELC m SLC m TECK m MIP-1γ	1C-50 1 nM 4 nM 2 nM 70 nM			

FIG. 4(b)

FIG. 5

CCXCKR	AIGCAGCAIC		ANGCCMCIA	GIGAACITA	GIGCAMATGC	20
5' upstream CCXCXR	TOAGAGAATT	TATTTAACTT	ATTTAAATTA	AATTIATAAA	TAACATCAAA	1.00
5' upstream CCXCKR	ATAAAAATA	AATTTAATTT	AAATAAACCA	AGTAATTTGC	TATTTICCTT	150
5' upstream CCXCKR	TTTATTCAAT			CGATTCACAA		200
5' upstream	GTAAAGATTA	TAACACTATT	TATTCTTTT	AGTTAAAATC	TAATTAAATT	250
5 upstream CCXCKR	TTCATATTT	AAAAATCATT	TTTACATAAA	AGTCTTCACT	TTTATTTAGG	300
i⊓ GCXCK& ∰ upstream	ATTTAATGAT	TAAGAAAATT	CTCCAGGGCA	TTATGTTTAT	TETCCTGTTC	350
Upstream CCXCKR	AAATCCAAGC			AAGCAAAGIT		400
2, nbscream	ATCTTGGGGT	CATATICCAA	TGTGGCTCCC	ATTAAAGCAT	TTCAAAGAGT	450
55 upstream ecxckR	GCTAGATTCA	GGCTCACATA	TGTTACAGCA	ACAGGCTATA	CTCTAGGGAA	500
Supstream CCXCKR	AGAACAAAAC	AGCTINATAG	AAACTGTGTG		TATTTAGACA	550 -
S. nbecreww	AATATCTATC	CTGTATTCTC	TTTGCCATCT			600 9
5' upstream CCXCXR	CANEAGAACE CAACAGAACE					649 58
5' upstream CCXCKR	CO ST ATEN	OFFICE CO.	mans a vence vences	ATCAA	REPLES AS A S A S A S A S A S A S A S A S A	685 108
5 'upstream CCXCKR	AGAGAN	CACAGGATAT		mile and a second		734 147
5' upstream CCXCKR	DAGE CED	TCATTGGACT	TGCAGGCAAT	TCCATGOTA	IGGCAATTTA	740 197
CCXCKR CCXCKR	TGCCTATTAC	aagaaacaga	GAACCAAAAC	AGATGIGIAC	ATCCTGAATT	74D 247
5' upstream CCXCRR	TGGCTGTAGC	AGATITACIC	CTTCTATTCA	CTCTGCCTT	TIGGGCTGTT	740 297
S' upstream	AATGCAGTTC	ATGGGTGGGT	TTTAGGGAAA	ATAATGTGCA	AAATAACTIC	740 347

% of Baseline Surface Expression

Fig. 6A

FLC

100 mM

1nB

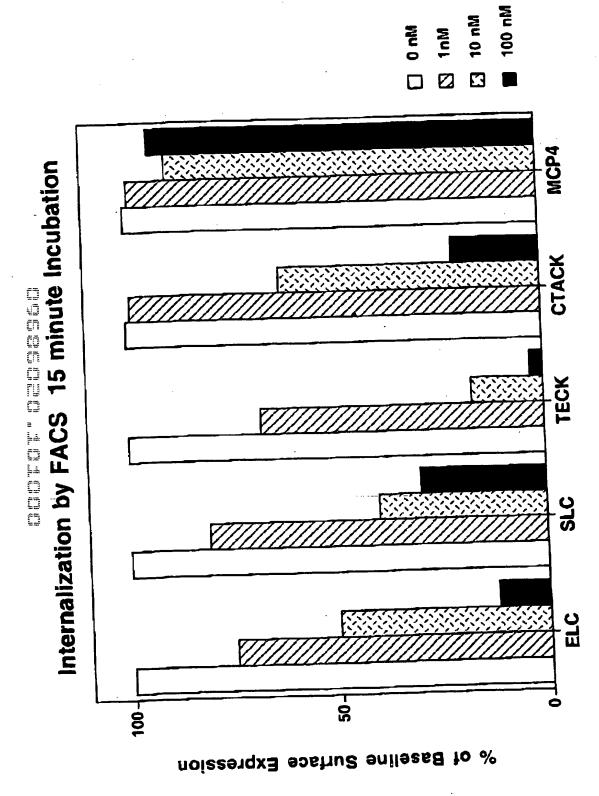


Fig. 6B